dbGaP Study Release Notes



Release Notes for NIGMS hTERT Tandem Repeats, phs002428.v1.p1

"Tandem DNA Repeats Activate hTERT Gene Transcription"

For any questions or comments, please contact: dbgap-help@ncbi.nlm.nih.gov.

June 9, 2021 Version 1 Data set release date

2021-06-09

Version 1 Data set release for NIGMS hTERT Tandem Repeats now available

This release includes phenotypes tables and tandem repeats. Please refer to the latest study configuration report for a detailed description of each download component.

Consent group 1 (c1): General Research Use (GRU)

Data Type	subjects	samples
Phenotype	209	418
Seq_Interactions	209	418

For a description of SAMPLE USE terms, please see:

https://www.ncbi.nlm.nih.gov/projects/gap/submission/GetSampleUseTypes.cgi

Molecular Data

- 1. Submitted molecular data are accessioned under phg001573.v1. Please see the folder marked as "sample-info" for mappings of subject, consent status, and sequence data files.
- 2. Sample-level sequences of tandem repeats are in original matrix format. The folder is named
 - "phg001573.v1.hTERT_TandemRepeats.original_submission_matrixfmt.target_markerset_gr c38.c1.GRU.tar.gz".

Authorized Access (Individual Level Data)

Individual level data are available for download through the dbGaP Authorized Access System upon approval of the Data Access Request (DAR):

• https://dbgap.ncbi.nlm.nih.gov/aa/wga.cgi?login=&page=login

Public FTP site (Summary Level Data Only)

All data tables and data dictionaries will be housed under one directory for ease of downloading. The data_dict filenames have an added study version number (phs#.v#) and deleted participant set number (p#) from the table accession (pht#.v#). The var_report filenames have an added study version number (phs#.v#). In the var_report files, variables contain version numbers (phv#.v#) and summaries were created for each consent group (c#). These FTP files are available at:

https://ftp.ncbi.nlm.nih.gov/dbgap/studies/phs002428/phs002428.v1.p1